

*Please cancel claim 2, without prejudice.*

*Please amend claim 3 as follows.*

B2 ~~3.~~ (Amended) Bacteria of claim 1, in which the lysE gene of *Corynebacterium glutamicum* encoding the lysine export carrier is over-expressed.

*Please cancel claims 4, 19, and 20, without prejudice*

*Please amend claims 22 and 23 as follows.*

B3 ~~22.~~ (Amended) An isolated DNA comprising the nucleotide sequence shown in SEQ ID No: 5.

~~23.~~ (Amended) An isolated DNA comprising the nucleotide sequence shown in SEQ ID No: 6.

*Please add following new claims 27-32.*

27. Bacteria of claim 1 further comprising an overexpressed lysC gene of *Corynebacterium glutamicum* encoding aspartate kinase.

B4 ~~28.~~ Bacteria of claim 27, wherein said over-expressed aspartate kinase is resistant to inhibition by lysine and/or threonine.

29. Bacteria of claim 1, wherein said dapA gene including a dapA promotor selected from the group consisting of: the dapA promotor comprising the MC20 mutation as set forth in SEQ ID NO:5 and the dapA promotor comprising the MA20 mutation as set forth in SEQ ID NO:6 is comprised in the aecD (amino ethyl cysteine degrading) gene of *Corynebacterium glutamicum*.

30. L-Lysine-producing bacteria of the species *Corynebacterium glutamicum* comprising an over-expressed pyc gene of *Corynebacterium glutamicum* encoding pyruvate carboxylase, an over-expressed dapA gene of *Corynebacterium glutamicum* encoding